

Sequence Listings

(1) GENERAL INFORMATION:

- 5 (i) INVENTORS: B. Mollet
R.D. Pridmore
M.C. Zwahlen
- 10 (ii) TITLE OF THE INVENTION: Novel strains of the Bacillus subtilis group
for food fermentation
- (iii) NUMBER OF SEQUENCES: 4
- 15 (iv) ADDRESS:
(A) Address: Winston & Strawn
(B) Street: 200 Park Avenue
(C) City: New York
(D) State: New York
(E) Country: USA
20 (F) Regional Code:
- (v) COMPUTER READABLE FORM:
(A) Medium type: Floppy Disk
25 (B) Computer: IBM PC Compatible
(C) System: PC-DOS/MS-DOS
(D) Software: Patentin Release Nr. 1, Version Nr. 1,25
- (vi) INFORMATION OF THE APPLICATION:
(A) Application number: to be obtained
30 (B) Application date: herewith
(C) Classification:
- (vii) PRIORITY DATES: none
- 35 (viii) Attorney/Agent Information
(A) Name: Allan A. Fanucci
(B) Registration number: 30,256
(C) Reference: 88265-4011
- 40 (ix) TELECOMMUNICATION:
(A) Telephone: 212-294-3311
(B) Telefax: 212-294-4700
- (2) INFORMATION FOR SEQ ID. NO. 1:
- 45 (i) SEQUENCE CHARACTERISTICS:
(A) length: 30 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
(D) topology: linear
- 50 (ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 1:

GCGGCGGATC CGCTGATGAT CTCCCAGCCC

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5 (3) INFORMATION FOR SEQ ID. NO. 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) length: 44 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
(D) topology: linear

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(ii) MOLECULE TYPE: DNA

(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 2:

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CTCAAATTCC ATTCCTCAT CAGGACATGC ATAGCGTATC ATCC

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(4) INFORMATION FOR SEQ ID. NO. 3:

(i) SEQUENCE CHARACTERISTICS:

- (A) length: 31 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
(D) Topology: linear

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(ii) MOLECULE TYPE: DNA

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(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 3:

GGGGTCGAAT TCCACGAGAT ATCTAACTGC C

31

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(5) INFORMATION FOR SEQ ID. NO. 4:

(i) SEQUENCE CHARACTERISTICS:

- (A) length: 44 nucleotides
(B) type: nucleic acid
(C) strandedness: single stranded
(D) Topology: linear

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(ii) MOLECULE TYPE: DNA

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(iii) SEQUENCE DESCRIPTION: SEQ ID. NO. 4:

GGATGATACG CTATGCATGT CCTGATGAGG AAATGGAATTTGAG

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